
Sequence Listing was accepted.

If you need help call the Patent Electronic Business Center at (866) 217-9197 (toll free).

Reviewer: Anne Corrigan

Timestamp: Thu Sep 27 15:01:55 EDT 2007

Validated By CRFValidator v 1.0.3

Application No: 10580542 Version No: 1.0

Input Set:

Output Set:

Started: 2007-09-27 14:48:38.334

Finished: 2007-09-27 14:48:39.197

Elapsed: 0 hr(s) 0 min(s) 0 sec(s) 863 ms

Total Warnings: 9

Total Errors: 0

No. of SeqIDs Defined: 15

Actual SeqID Count: 15

Error code		Error Description	า							
W	213	Artificial o	r Unknown	found	in	<213>	in	SEQ	ID	(5)
W	213	Artificial o	r Unknown	found	in	<213>	in	SEQ	ID	(8)
W	213	Artificial o	r Unknown	found	in	<213>	in	SEQ	ID	(9)
W	213	Artificial o	r Unknown	found	in	<213>	in	SEQ	ID	(10)
W	213	Artificial o	r Unknown	found	in	<213>	in	SEQ	ID	(11)
W	213	Artificial o	r Unknown	found	in	<213>	in	SEQ	ID	(12)
W	213	Artificial o	r Unknown	found	in	<213>	in	SEQ	ID	(13)
W	213	Artificial o	r Unknown	found	in	<213>	in	SEQ	ID	(14)
W	213	Artificial o	r Unknown	found	in	<213>	in	SEQ	ID	(15)

SEQUENCE LISTING

<110>	0> Wallach, David Ramakrishnan, Parameswaran Shmushkovich, Taisia Wang, Wangxia													
<120>	METHODS OF REGULATING AN IMMUNE RESPONSE													
<130>	27083													
<140>	10580542													
<141>	2007-09-27													
<160>	15													
<170>	> PatentIn version 3.2													
<210>	1													
<211>	2844													
<212>	DNA													
<213>	Homo sapiens													
<400>	1													
atggcag	ıtga tggaaatggc c	tgcccaggt	gcccctggct	cagcagtggg	gcagcagaag	60								
gaactco	cca agccaaagga g	aagacgccg	ccactgggga	agaaacagag	ctccgtctac	120								
aagctto	gagg ccgtggagaa g	agccctgtg	ttctgcggaa	agtgggagat	cctgaatgac	180								
gtgatta	icca agggcacagc c	aaggaaggc	tccgaggcag	ggccagctgc	catctctatc	240								
atcgccc	agg ctgagtgtga g	aatagccaa	gagttcagcc	ccaccttttc	agaacgcatt	300								
ttcatco	yctg ggtccaaaca g	tacagccag	tccgagagtc	ttgatcagat	ccccaacaat	360								
gtggcc	atg ctacagaggg c	aaaatggcc	cgtgtgtgtt	ggaagggaaa	gcgtcgcagc	420								
aaagcco	gga agaaacggaa g	aagaagagc	tcaaagtccc	tggctcatgc	aggagtggcc	480								
ttggcca	aac ccctccccag g	acccctgag	caggagagct	gcaccatccc	agtgcaggag	540								
gatgagt	ctc cactcggcgc c	ccatatgtt	agaaacaccc	cgcagttcac	caagcctctg	600								
aaggaad	cag gccttgggca a	ctctgtttt.	aagcagcttg	gcgagggcct	acggccggct	660								
ctgcct	gat cagaactcca c	aaactgatc	agccccttgc	aatgtctgaa	ccacgtgtgg	720								
						= 4 -								
aaactgo	cacc acccccagga c	ggaggcccc	ctgcccctgc	ccacgcaccc	cttcccctat	780								
						2.4.5								
agcagac	tgc ctcatccctt c	CCALLCCAC	cererecage	cciggaaacc	Leacectetg	840								

gagtccttcc tgggcaaact ggcctgtgta gacagccaga aacccttgcc tgacccacac

ctgagcaaac tggcctgtgt agacagtcca aagcccctgc ctggcccaca cctggagccc

900

960

agctgcctgt ct	tegtggtge	ccatgagaag	ttttctgtgg	aggaatacct	agtgcatgct	1020
ctgcaaggca go	cgtgagctc	aagccaggcc	cacagcctga	ccagcctggc	caagacctgg	1080
gcagcacggg go	ctccagatc	ccgggagccc	agccccaaaa	ctgaggacaa	cgagggtgtc	1140
ctgctcactg ac	gaaactcaa	gccagtggat	tatgagtacc	gagaagaagt	ccactgggcc	1200
acgcaccagc to	ccgcctggg	cagaggctcc	ttcggagagg	tgcacaggat	ggaggacaag	1260
cagactggct to	ccagtgcgc	tgtcaaaaag	gtgcggctgg	aagtatttcg	ggcagaggag	1320
ctgatggcat gt	tgcaggatt	gacctcaccc	agaattgtcc	ctttgtatgg	agctgtgaga	1380
gaagggcctt go	ggtcaacat	cttcatggag	ctgctggaag	gtggctccct	gggccagctg	1440
gtcaaggagc ag	gggctgtct	cccagaggac	cgggccctgt	actacctggg	ccaggccctg	1500
gagggtctgg aa	atacctcca	ctcacgaagg	attctgcatg	gggacgtcaa	agctgacaac	1560
gtgctcctgt co	cagcgatgg	gagccacgca	gecetetgtg	actttggcca	tgctgtgtgt	1620
cttcaacctg at	tggcctggg	aaagtccttg	ctcacagggg	actacatccc	tggcacagag	1680
acccacatgg ct	tccggaggt	ggtgctgggc	aggagctgcg	acgccaaggt	ggatgtctgg	1740
agcagctgct gt	tatgatgct	gcacatgctc	aacggctgcc	acccctggac	tcagttcttc	1800
cgagggccgc to	ctgcctcaa	gattgccagc	gageeteege	ctgtgaggga	gatcccaccc	1860
teetgegeee et	tctcacagc	ccaggccatc	caagaggggc	tgaggaaaga	gcccatccac	1920
cgcgtgtctg ca	agcggagct	gggagggaag	gtgaaccggg	cactacagca	agtgggaggt	1980
ctgaagagcc ct	ttggagggg	agaatataaa	gaaccaagac	atccaccgcc	aaatcaagcc	2040
aattaccacc ac	gaccctcca	tgcccagccg	agagagcttt	cgccaagggc	cccagggccc	2100
eggeeagetg ac	ggagacaac	aggcagagcc	cctaagctcc	agcctcctct	cccaccagag	2160
cccccagagc ca	aaacaagtc	tectecettg	actttgagca	aggaggagtc	tgggatgtgg	2220
gaacccttac ct	tctgtcctc	cctggagcca	gcccctgcca	gaaaccccag	ctcaccagag	2280
cggaaagcaa co	cgtcccgga	gcaggaactg	cagcagctgg	aaatagaatt	attcctcaac	2340
agcctgtccc ac	gccattttc	tctggaggag	caggagcaaa	ttctctcgtg	cctcagcatc	2400
gacageetet ee	cctgtcgga	tgacagtgag	aagaacccat	caaaggcctc	tcaaagctcg	2460
egggacacee to	gageteagg	cgtacactcc	tggagcagcc	aggccgaggc	tcgaagctcc	2520
agctggaaca to	ggtgctggc	ccgggggcgg	cccaccgaca	ccccaagcta	tttcaatggt	2580
gtgaaagtcc aa	aatacagtc	tcttaatggt	gaacacctgc	acatccggga	gttccaccgg	2640
gtcaaagtgg ga	agacatcgc	cactggcatc	agcagccaga	teccagetge	agccttcagc	2700

ttggtcacca aagacgggca gcctgttcgc tacgacatgg aggtgccaga ctcgggcatc	2760												
gacctgcagt gcacactggc ccctgatggc agcttcgcct ggagctggag ggtcaagcat	2820												
ggccagctgg agaacaggcc ctaa	2844												
<210> 2 <211> 947 <212> PRT <213> Homo sapiens <400> 2													
Met Ala Val Met Glu Met Ala Cys Pro Gly Ala Pro Gly Ser Ala Val 1 5 10 15													
Gly Gln Gln Lys Glu Leu Pro Lys Pro Lys Glu Lys Thr Pro Pro Leu 20 25 30													
Gly Lys Lys Gln Ser Ser Val Tyr Lys Leu Glu Ala Val Glu Lys Ser 35 40 45													
Pro Val Phe Cys Gly Lys Trp Glu Ile Leu Asn Asp Val Ile Thr Lys 50 55 60													
Gly Thr Ala Lys Glu Gly Ser Glu Ala Gly Pro Ala Ala Ile Ser Ile 65 70 75 80													
Ile Ala Gln Ala Glu Cys Glu Asn Ser Gln Glu Phe Ser Pro Thr Phe 85 90 95													
Ser Glu Arg Ile Phe Ile Ala Gly Ser Lys Gln Tyr Ser Gln Ser Glu 100 105 110													
Ser Leu Asp Gln Ile Pro Asn Asn Val Ala His Ala Thr Glu Gly Lys 115 120 125													
Met Ala Arg Val Cys Trp Lys Gly Lys Arg Arg Ser Lys Ala Arg Lys 130 135 140													
Lys Arg Lys Lys Ser Ser Lys Ser Leu Ala His Ala Gly Val Ala 145 150 155 160													

Leu Ala Lys Pro Leu Pro Arg Thr Pro Glu Gln Glu Ser Cys Thr Ile

Pro	Val	Gln	Glu 180	Asp	Glu	Ser	Pro	Leu 185	Gly	Ala	Pro	Tyr	Val 190	Arg	Asn
Thr	Pro	Gln 195	Phe	Thr	Lys	Pro	Leu 200	Lys	Glu	Pro	Gly	Leu 205	Gly	Gln	Leu
Cys	Phe 210	Lys	Gln	Leu	Gly	Glu 215	Gly	Leu	Arg	Pro	Ala 220	Leu	Pro	Arg	Ser
Glu 225	Leu	His	Lys	Leu	Ile 230	Ser	Pro	Leu	Gln	Cys 235	Leu	Asn	His	Val	Trp 240
Lys	Leu	His	His	Pro 245	Gln	Asp	Gly	Gly	Pro 250	Leu	Pro	Leu	Pro	Thr 255	His
Pro	Phe	Pro	Tyr 260	Ser	Arg	Leu	Pro	His 265	Pro	Phe	Pro	Phe	His 270	Pro	Leu
Gln	Pro	Trp 275	Lys	Pro	His	Pro	Leu 280	Glu	Ser	Phe	Leu	Gly 285	Lys	Leu	Ala
Суз	Val 290	Asp	Ser	Gln	Lys	Pro 295	Leu	Pro	Asp	Pro	His 300	Leu	Ser	Lys	Leu
Ala 305	Cys	Val	Asp	Ser	Pro 310	Lys	Pro	Leu	Pro	Gly 315	Pro	His	Leu	Glu	Pro 320
Ser	Cys	Leu	Ser	Arg 325	Gly	Ala	His	Glu	Lys 330	Phe	Ser	Val	Glu	Glu 335	Tyr
Leu	Val	His	Ala 340	Leu	Gln	Gly	Ser	Val 345	Ser	Ser	Ser	Gln	Ala 350	His	Ser
Leu	Thr	Ser 355	Leu	Ala	Lys	Thr	Trp 360	Ala	Ala	Arg	Gly	Ser 365	Arg	Ser	Arg
Glu	Pro 370	Ser	Pro	Lys	Thr	Glu 375	Asp	Asn	Glu	Gly	Val 380	Leu	Leu	Thr	Glu
Lys 385	Leu	Lys	Pro	Val	Asp 390	Tyr	Glu	Tyr	Arg	Glu 395	Glu	Val	His	Trp	Ala 400

Thr	His	Gln	Leu	Arg 405	Leu	Gly	Arg	Gly	Ser 410	Phe	Gly	Glu	Val	His 415	Arg
Met	Glu	Asp	Lys 420	Gln	Thr	Gly	Phe	Gln 425	Суз	Ala	Val	Lys	Lys 430	Val	Arg
Leu	Glu	Val 435	Phe	Arg	Ala	Glu	Glu 440	Leu	Met	Ala	Суз	Ala 445	Gly	Leu	Thr
Ser	Pro 450	Arg	Ile	Val	Pro	Leu 455	Tyr	Gly	Ala	Val	Arg 460	Glu	Gly	Pro	Trp
Val 465	Asn	Ile	Phe	Met	Glu 470	Leu	Leu	Glu	Gly	Gly 475	Ser	Leu	Gly	Gln	Leu 480
Val	Lys	Glu	Gln	Gly 485	Суѕ	Leu	Pro	Glu	Asp 490	Arg	Ala	Leu	Tyr	Tyr 495	Leu
Gly	Gln	Ala	Leu 500	Glu	Gly	Leu	Glu	Tyr 505	Leu	His	Ser	Arg	Arg 510	Ile	Leu
His	Gly	Asp 515	Val	Lys	Ala	Asp	Asn 520	Val	Leu	Leu	Ser	Ser 525	Asp	Gly	Ser
His	Ala 530	Ala	Leu	Cys	Asp	Phe 535	Gly	His	Ala	Val	Cys 540	Leu	Gln	Pro	Asp
Gly 545	Leu	Gly	Lys	Ser	Leu 550	Leu	Thr	Gly	Asp	Tyr 555	Ile	Pro	Gly	Thr	Glu 560
Thr	His	Met	Ala	Pro 565	Glu	Val	Val	Leu	Gly 570	Arg	Ser	Суз	Asp	Ala 575	Lys
Val	Asp	Val	Trp 580	Ser	Ser	Суѕ	Суз	Met 585	Met	Leu	His	Met	Leu 590	Asn	Gly
Cys	His	Pro 595	Trp	Thr	Gln	Phe	Phe 600	Arg	Gly	Pro	Leu	Cys 605	Leu	Lys	Ile
Ala	Ser 610	Glu	Pro	Pro	Pro	Val 615	Arg	Glu	Ile	Pro	Pro 620	Ser	Суз	Ala	Pro

Leu 625	Thr	Ala	Gln	Ala	Ile 630	Gln	Glu	Gly	Leu	Arg 635	Lys	Glu	Pro	Ile	His 640
Arg	Val	Ser	Ala	Ala 645	Glu	Leu	Gly	Gly	Lys 650	Val	Asn	Arg	Ala	Leu 655	Gln
Gln	Val	Gly	Gly 660	Leu	Lys	Ser	Pro	Trp 665	Arg	Gly	Glu	Tyr	Lys 670	Glu	Pro
Arg	His	Pro 675	Pro	Pro	Asn	Gln	Ala 680	Asn	Tyr	His	Gln	Thr 685	Leu	His	Ala
Gln	Pro 690	Arg	Glu	Leu	Ser	Pro 695	Arg	Ala	Pro	Gly	Pro 700	Arg	Pro	Ala	Glu
Glu 705	Thr	Thr	Gly	Arg	Ala 710	Pro	Lys	Leu	Gln	Pro 715	Pro	Leu	Pro	Pro	Glu 720
Pro	Pro	Glu	Pro	Asn 725	Lys	Ser	Pro	Pro	14 Table 14 Table 15 Table 16	Thr	Leu	Ser	Lys	Glu 735	Glu
			740					745					Pro 750		
Ala	_	755					760		-			765			
	770					775					780		Leu		
785					790					795		-	Leu		800
Asp				805		-	-		810	-				815	
Ser	Gln	Ser	Ser 820	Arg	Asp	Thr	Leu	Ser 825	Ser	Gly	Val	His	Ser 830	Trp	Ser
Ser	Gln	Ala 835	Glu	Ala	Arg	Ser	Ser 840	Ser	Trp	Asn	Met	Val 845	Leu	Ala	Arg

Gly Arg Pro Thr Asp Thr Pro Ser Tyr Phe Asn Gly Val Lys Val Gln

850 855 860

Ile Gln Ser Leu Asn Gly Glu His Leu His Ile Arg Glu Phe His Arg 865 870 875 880

Val Lys Val Gly Asp Ile Ala Thr Gly Ile Ser Ser Gln Ile Pro Ala 885 890 895

Ala Ala Phe Ser Leu Val Thr Lys Asp Gly Gln Pro Val Arg Tyr Asp 900 905 910

Met Glu Val Pro Asp Ser Gly Ile Asp Leu Gln Cys Thr Leu Ala Pro 915 920 925

Asp Gly Ser Phe Ala Trp Ser Trp Arg Val Lys His Gly Gln Leu Glu 930 935 940

Asn Arg Pro 945

<210> 3

<211> 175

<212> PRT

<213> Homo sapiens

<400> 3

Met Pro Lys Arg Ser Cys Pro Phe Ala Asp Val Ala Pro Leu Gln Leu 1 5 10 15

Lys Val Arg Val Ser Gln Arg Glu Leu Ser Arg Gly Val Cys Ala Glu
20 25 30

Arg Tyr Ser Gln Glu Val Phe Glu Lys Thr Lys Arg Leu Leu Phe Leu 35 40 45

Gly Ala Gln Ala Tyr Leu Asp His Val Trp Asp Glu Gly Cys Ala Val 50 55 60

Val His Leu Pro Glu Ser Pro Lys Pro Gly Pro Thr Gly Ala Pro Arg 65 70 75 80

Ala Ala Arg Gly Gln Met Leu Ile Gly Pro Asp Gly Arg Leu Ile Arg 85 90 95 Ser Leu Gly Gln Ala Ser Glu Ala Asp Pro Ser Gly Val Ala Ser Ile 100 105 110

Ala Cys Ser Ser Cys Val Arg Ala Val Asp Gly Lys Ala Val Cys Gly
115 120 125

Gln Cys Glu Arg Ala Leu Cys Gly Gln Cys Val Arg Thr Cys Trp Gly 130 135 140

Cys Gly Ser Val Ala Cys Thr Leu Cys Gly Leu Val Asp Cys Ser Asp 145 150 155 160

Met Tyr Glu Lys Val Leu Cys Thr Ser Cys Ala Met Phe Glu Thr $165 \hspace{1.5cm} 170 \hspace{1.5cm} 175$

<210> 4

<211> 110

<212> PRT

<213> Homo sapiens

<400> 4

Met Pro Lys Arg Ser Cys Pro Phe Ala Asp Val Ala Pro Leu Gln Leu 1 5 10 15

Lys Val Arg Val Ser Gln Arg Glu Leu Ser Arg Gly Val Cys Ala Glu 20 25 30

Arg Tyr Ser Gln Glu Val Phe Asp Pro Ser Gly Val Ala Ser Ile Ala 35 4045

Cys Ser Ser Cys Val Arg Ala Val Asp Gly Lys Ala Val Cys Gly Gln 50 55 60

Cys Glu Arg Ala Leu Cys Gly Gln Cys Val Arg Thr Cys Trp Gly Cys 65 70 75 80

Gly Ser Val Ala Cys Thr Leu Cys Gly Leu Val Asp Cys Ser Asp Met 85 90 95

Tyr Glu Lys Val Leu Cys Thr Ser Cys Ala Met Phe Glu Thr 100 105 110

```
<212> PRT
<213> Artificial sequence
<220>
<223> Myc tag
<400> 5
Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
<210> 6
<211> 528
<212> DNA
<213> Homo sapiens
<400> 6
atgcccaage ggagetgeee ettegeggae gtggeeeege tacageteaa ggteegegtg
agccagaggg agttgagccg cggcgtgtgc gccgagcgct actcgcagga ggtcttcgag
                                                                    120
aagaccaagc gactcctgtt cctcggggcc caggcctacc tggaccacgt gtgggatgaa
                                                                    180
ggctgtgccg tcgttcacct gccagagtcc ccaaagcctg gccctacagg ggccccgagg
                                                                    240
                                                                    300
gctgcacgtg ggcagatgct gattggacca gacggccgcc tgatcaggag ccttgggcag
gcctccgaag ctgacccatc tggggtagcg tccattgcct gttcctcatg cgtgcgagcc
                                                                    360
gtggatggga aggcggtctg cggtcagtgt gagcgagccc tgtgcgggca gtgtgtgcgc
                                                                    420
                                                                    480
acctgctggg gctgcggctc cgtggcctgt accctgtgtg gcctcgtgga ctgcagtgac
                                                                    528
atgtacgaga aagtgctgtg caccagctgt gccatgttcg agacctga
<210> 7
<211> 333
<212> DNA
<213> Homo sapiens
<400> 7
atgcccaage ggagetgeee ettegeggae gtggeeeege tacageteaa ggteegegtg
                                                                     60
agccagaggg agttgagccg cggcgtgtgc gccgagcgct actcgcagga ggtcttcgac
                                                                    120
                                                                    180
ccatctgggg tagcgtccat tgcctgttcc tcatgcgtgc gagccgtgga tgggaaggcg
gtctgcggtc agtgtgagcg agccctgtgc gggcagtgtg tgcgcacctg ctggggctgc
                                                                  240
ggctccgtgg cctgtaccct gtgtggcctc gtggactgca gtgacatgta cgagaaagtg
                                                                    300
                                                                    333
ctgtgcacca gctgtgccat gttcgagacc tga
```

<211> 10

```
<211> 34
<212> DNA
<213> Artificial sequence
<220>
<223> Single strand DNA oligonucleotide
ccaagctatt tcaatcgtgt gaaagtccaa atac
                                                                      34
<210> 9
<211> 34
<212> DNA
<213> Artificial sequence
<220>
<223> Single strand DNA oligonucleotide
<400> 9
gtatttggac tttcacacga ttgaaatagc ttgg
                                                                      34
<210> 10
<211> 16
<212> PRT
<213> Artificial sequence
<220>
<223> A peptide corresponding to a sequence within the NIK kinase
       domain
<400> 10
Arg Leu Gly Arg Gly Ser Phe Gly Glu Val His Arg Met Glu Asp Lys
                                   10
                                                       15
<210> 11
<211> 42
<212> DNA
<213> Artificial sequence
<220>
<223> Single strand DNA oligonucleotide
<400> 11
                                                                      42
gagggtctgg aatacctaca ttcccgcagg attctgcatg gg
<210> 12
<211> 42
<212> DNA
<213> Artificial sequence
```

<210> 8